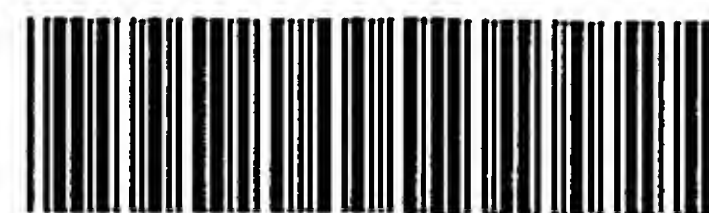


RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/594,117
Source: JFWP
Date Processed by STIC: 10/06/2006

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IFWP

RAW SEQUENCE LISTING

DATE: 10/06/2006

PATENT APPLICATION: US/10/594,117

TIME: 11:10:29

Input Set : A:\004974.01219 sequence listing.txt

Output Set: N:\CRF4\10062006\J594117.raw

3 <110> APPLICANT: Golz, Stefan
 4 Bruggemeier, Ulf
 5 Geerts, Andreas
 6 Summer, Holger
 8 <120> TITLE OF INVENTION: Diagnostics and Therapeutics for Diseases Associated with
 Protein
 9 Kinase, cGMP-Dependent, Type I (PRKG1)
 11 <130> FILE REFERENCE: 004974.01219
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/594,117
 C--> 12 <141> CURRENT FILING DATE: 2006-09-25
 12 <150> PRIOR APPLICATION NUMBER: PCT/EP2005/02531
 13 <151> PRIOR FILING DATE: 2005-03-10
 15 <150> PRIOR APPLICATION NUMBER: EP 04007085.6
 16 <151> PRIOR FILING DATE: 2004-03-24
 18 <160> NUMBER OF SEQ ID NOS: 5
 20 <170> SOFTWARE: PatentIn version 3.2
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 3740
 24 <212> TYPE: DNA
 25 <213> ORGANISM: Homo sapiens
 27 <400> SEQUENCE: 1
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 49 gtttgcaatg aagattctca agaaacgtca cattgtggac acaagacagc aggagcacat 1320
 50 ccgctcagag aagcagatca tgcagggggc tcattccgat ttcatagtga gactgtacag 1380

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53 atgtgtggta gaagcttttg cctatctgca ttccaaagga atcatttaca gggacctcaa 1560
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83 accaaaaaac aacaaacaaa caaaaaacaa gaatgaaaaa cagaaataaa agaagtagaa 3360
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3740

92 <210> SEQ ID NO: 2

93 <211> LENGTH: 686

94 <212> TYPE: PRT

95 <213> ORGANISM: Homo sapiens

97 <400> SEQUENCE: 2

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99 1 5 10 15

100 Glu Leu Arg Gln Arg Asp Ala Leu Ile Asp Glu Leu Glu Leu Glu Leu

101 20 25 30

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104	Tyr	Arg	Ser	Val	Ile	Arg	Pro	Ala	Thr	Gln	Gln	Ala	Gln	Lys	Gln	Ser
105		50					55					60				
106	Ala	Ser	Thr	Leu	Gln	Gly	Glu	Pro	Arg	Thr	Lys	Arg	Gln	Ala	Ile	Ser
107	65					70				75						80
108	Ala	Glu	Pro	Thr	Ala	Phe	Asp	Ile	Gln	Asp	Leu	Ser	His	Val	Thr	Leu
109				85						90					95	
110	Pro	Phe	Tyr	Pro	Lys	Ser	Pro	Gln	Ser	Lys	Asp	Leu	Ile	Lys	Glu	Ala
111				100					105					110		
112	Ile	Leu	Asp	Asn	Asp	Phe	Met	Lys	Asn	Leu	Glu	Leu	Ser	Gln	Ile	Gln
113			115					120					125			
114	Glu	Ile	Val	Asp	Cys	Met	Tyr	Pro	Val	Glu	Tyr	Gly	Lys	Asp	Ser	Cys
115		130					135					140				
116	Ile	Ile	Lys	Glu	Gly	Asp	Val	Gly	Ser	Leu	Val	Tyr	Val	Met	Glu	Asp
117	145					150					155				160	
118	Gly	Lys	Val	Glu	Val	Thr	Lys	Glu	Gly	Val	Lys	Leu	Cys	Thr	Met	Gly
119					165					170					175	
120	Pro	Gly	Lys	Val	Phe	Gly	Glu	Leu	Ala	Ile	Leu	Tyr	Asn	Cys	Thr	Arg
121				180					185					190		
122	Thr	Ala	Thr	Val	Lys	Thr	Leu	Val	Asn	Val	Lys	Leu	Trp	Ala	Ile	Asp
123			195						200					205		
124	Arg	Gln	Cys	Phe	Gln	Thr	Ile	Met	Met	Arg	Thr	Gly	Leu	Ile	Lys	His
125		210					215					220				
126	Thr	Glu	Tyr	Met	Glu	Phe	Leu	Lys	Ser	Val	Pro	Thr	Phe	Gln	Ser	Leu
127	225					230					235				240	
128	Pro	Glu	Glu	Ile	Leu	Ser	Lys	Leu	Ala	Asp	Val	Leu	Glu	Glu	Thr	His
129				245						250					255	
130	Tyr	Glu	Asn	Gly	Glu	Tyr	Ile	Ile	Arg	Gln	Gly	Ala	Arg	Gly	Asp	Thr
131			260						265					270		
132	Phe	Phe	Ile	Ile	Ser	Lys	Gly	Thr	Val	Asn	Val	Thr	Arg	Glu	Asp	Ser
133			275					280					285			
134	Pro	Ser	Glu	Asp	Pro	Val	Phe	Leu	Arg	Thr	Leu	Gly	Lys	Gly	Asp	Trp
135		290					295					300				
136	Phe	Gly	Glu	Lys	Ala	Leu	Gln	Gly	Glu	Asp	Val	Arg	Thr	Ala	Asn	Val
137	305					310					315				320	
138	Ile	Ala	Ala	Glu	Ala	Val	Thr	Cys	Leu	Val	Ile	Asp	Arg	Asp	Ser	Phe
139				325						330					335	
140	Lys	His	Leu	Ile	Gly	Gly	Leu	Asp	Asp	Val	Ser	Asn	Lys	Ala	Tyr	Glu
141			340					345						350		
142	Asp	Ala	Glu	Ala	Lys	Ala	Lys	Tyr	Glu	Ala	Glu	Ala	Ala	Phe	Phe	Ala
143			355					360					365			
144	Asn	Leu	Lys	Leu	Ser	Asp	Phe	Asn	Ile	Ile	Asp	Thr	Leu	Gly	Val	Gly
145		370					375					380				
146	Gly	Phe	Gly	Arg	Val	Glu	Leu	Val	Gln	Leu	Lys	Ser	Glu	Glu	Ser	Lys
147	385					390					395				400	
148	Thr	Phe	Ala	Met	Lys	Ile	Leu	Lys	Lys	Arg	His	Ile	Val	Asp	Thr	Arg
149				405						410					415	
150	Gln	Gln	Glu	His	Ile	Arg	Ser	Glu	Lys	Gln	Ile	Met	Gln	Gly	Ala	His

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DATE: 10/06/2006

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Output Set: N:\CRF4\10062006\J594117.raw

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153          435          440          445
154 Leu Tyr Met Leu Met Glu Ala Cys Leu Gly Gly Glu Leu Trp Thr Ile
155          450          455          460
156 Leu Arg Asp Arg Gly Ser Phe Glu Asp Ser Thr Thr Arg Phe Tyr Thr
157 465          470          475          480
158 Ala Cys Val Val Glu Ala Phe Ala Tyr Leu His Ser Lys Gly Ile Ile
159          485          490          495
160 Tyr Arg Asp Leu Lys Pro Glu Asn Leu Ile Leu Asp His Arg Gly Tyr
161          500          505          510
162 Ala Lys Leu Val Asp Phe Gly Phe Ala Lys Lys Ile Gly Phe Gly Lys
163          515          520          525
164 Lys Thr Trp Thr Phe Cys Gly Thr Pro Glu Tyr Val Ala Pro Glu Ile
165          530          535          540
166 Ile Leu Asn Lys Gly His Asp Ile Ser Ala Asp Tyr Trp Ser Leu Gly
167 545          550          555          560
168 Ile Leu Met Tyr Glu Leu Leu Thr Gly Ser Pro Pro Phe Ser Gly Pro
169          565          570          575
170 Asp Pro Met Lys Thr Tyr Asn Ile Ile Leu Arg Gly Ile Asp Met Ile
171          580          585          590
172 Glu Phe Pro Lys Lys Ile Ala Lys Asn Ala Ala Asn Leu Ile Lys Lys
173          595          600          605
174 Leu Cys Arg Asp Asn Pro Ser Glu Arg Leu Gly Asn Leu Lys Asn Gly
175          610          615          620
176 Val Lys Asp Ile Gln Lys His Lys Trp Phe Glu Gly Phe Asn Trp Glu
177 625          630          635          640
178 Gly Leu Arg Lys Gly Thr Leu Thr Pro Pro Ile Ile Pro Ser Val Ala
179          645          650          655
180 Ser Pro Thr Asp Thr Ser Asn Phe Asp Ser Phe Pro Glu Asp Asn Asp
181          660          665          670
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185 <210> SEQ ID NO: 3

186 <211> LENGTH: 20

187 <212> TYPE: DNA

188 <213> ORGANISM: artificial sequence

190 <220> FEATURE:

191 <223> OTHER INFORMATION: forward primer

193 <400> SEQUENCE: 3

194 agccgactac tggtcactgg

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196 <210> SEQ ID NO: 4

197 <211> LENGTH: 20

198 <212> TYPE: DNA

199 <213> ORGANISM: artificial sequence

201 <220> FEATURE:

202 <223> OTHER INFORMATION: reverse primer

204 <400> SEQUENCE: 4

205 gatctgggccc tgagaaaggt

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208 <211> LENGTH: 25
209 <212> TYPE: DNA
210 <213> ORGANISM: artificial sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: probe
215 <400> SEQUENCE: 5
216 tgtatgaact cctgactggc agccc

25

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/594,117

DATE: 10/06/2006

TIME: 11:10:30

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L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date